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<110> Shaughnessy, John D.
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 <223> Mouse Evi27 protein

<400> 7

Met	Leu	Leu	Val	Leu	Leu	Ile	Leu	Ala	Ala	Ser	Cys	Arg	Ser	Ala	
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Leu	Pro	Arg	Glu	Pro	Thr	Ile	Gln	Cys	Gly	Ser	Glu	Thr	Gly	Pro	
				20					25					30	
Ser	Pro	Glu	Trp	Met	Val	Gln	His	Thr	Leu	Thr	Pro	Gly	Asp	Leu	
				35					40					45	
Arg	Asp	Leu	Gln	Val	Glu	Leu	Val	Lys	Thr	Ser	Val	Ala	Ala	Glu	
				50					55					60	
Glu	Phe	Ser	Ile	Leu	Met	Asn	Ile	Ser	Trp	Ile	Leu	Arg	Ala	Asp	
				65					70					75	
Ala	Ser	Ile	Arg	Leu	Leu	Lys	Ala	Thr	Lys	Ile	Cys	Val	Ser	Gly	
				80					85					90	
Lys	Asn	Asn	Met	Asn	Ser	Tyr	Ser	Cys	Val	Arg	Cys	Asn	Tyr	Thr	
				95					100					105	
Glu	Ala	Phe	Gln	Ser	Gln	Thr	Arg	Pro	Ser	Gly	Gly	Lys	Trp	Thr	
				110					115					120	
Phe	Ser	Tyr	Val	Gly	Phe	Pro	Val	Glu	Leu	Ser	Thr	Leu	Tyr	Leu	
				125					130					135	
Ile	Ser	Ala	His	Asn	Ile	Pro	Asn	Ala	Asn	Met	Asn	Glu	Asp	Ser	
				140					145					150	
Pro	Ser	Leu	Ser	Val	Asn	Phe	Thr	Ser	Pro	Gly	Cys	Leu	Asn	His	
				155					160					165	
Val	Met	Lys	Tyr	Lys	Lys	Gln	Cys	Thr	Glu	Ala	Gly	Ser	Leu	Trp	
				170					175					180	

[illegible]

<210>	8
<211>	289
<212>	PRT
<213>	Unknown
<220>	

<221> PEPTIDE
 <223> IL-17 receptor protein

<400> 9

Met	Gly	Ala	Ala	Arg	Ser	Pro	Pro	Ser	Ala	Val	Pro	Gly	Pro	Leu	
				5					10					15	
Leu	Gly	Leu	Leu	Leu	Leu	Leu	Leu	Gly	Val	Leu	Ala	Pro	Gly	Gly	
				20					25					30	
Ala	Ser	Leu	Arg	Leu	Leu	Asp	His	Arg	Ala	Leu	Val	Cys	Ser	Gln	
				35					40					45	
Pro	Gly	Leu	Asn	Cys	Thr	Val	Lys	Asn	Ser	Thr	Cys	Leu	Asp	Asp	
				50					55					60	
Ser	Trp	Ile	His	Pro	Arg	Asn	Leu	Thr	Pro	Ser	Ser	Pro	Lys	Asp	
				65					70					75	
Leu	Gln	Ile	Gln	Leu	His	Phe	Ala	His	Thr	Gln	Gln	Gly	Asp	Leu	
				80					85					90	
Phe	Pro	Val	Ala	His	Ile	Glu	Trp	Thr	Leu	Gln	Thr	Asp	Ala	Ser	
				95					100					105	
Ile	Leu	Tyr	Leu	Glu	Gly	Ala	Glu	Leu	Ser	Val	Leu	Gln	Leu	Asn	
				110					115					120	
Thr	Asn	Glu	Arg	Leu	Cys	Val	Arg	Phe	Glu	Phe	Leu	Ser	Lys	Leu	
				125					130					135	
Arg	His	His	His	Arg	Arg	Trp	Arg	Phe	Thr	Phe	Ser	His	Phe	Val	
				140					145					150	
Val	Asp	Pro	Asp	Gln	Glu	Tyr	Glu	Val	Thr	Val	His	His	Leu	Pro	
				155					160					165	
Lys	Pro	Ile	Pro	Asp	Gly	Asp	Pro	Asn	His	Gln	Ser	Lys	Asn	Phe	
				170					175					180	
Leu	Val	Pro	Asp	Cys	Glu	His	Ala	Arg	Met	Lys	Val	Thr	Thr	Pro	
				185					190					195	
Cys	Met	Ser	Ser	Gly	Ser	Leu	Trp	Asp	Pro	Asn	Ile	Thr	Val	Glu	
				200					205					210	
Thr	Leu	Glu	Ala	His	Gln	Leu	Arg	Val	Ser	Phe	Thr	Leu	Trp	Asn	
				215					220					225	
Glu	Ser	Thr	His	Tyr	Gln	Ile	Leu	Leu	Thr	Ser	Phe	Pro	His	Met	
				230					235					240	
Glu	Asn	His	Ser	Cys	Phe	Glu	His	Met	His	His	Ile	Pro	Ala	Pro	
				245					250					255	
Arg	Pro	Glu	Glu	Phe	His	Gln	Arg	Ser	Asn	Val	Thr	Leu	Thr	Leu	
				260					265					270	
Arg	Asn	Leu	Lys	Gly	Cys	Cys	Arg	His	Gln	Val	Gln	Ile	Gln	Pro	
				275					280					285	
Phe	Phe	Ser	Ser	Cys	Leu	Asn	Asp	Cys	Leu	Arg	His	Ser	Ala	Thr	
				290					295					300	
Val	Ser	Cys	Pro	Glu	Met	Pro	Asp	Thr	Pro	Glu	Pro	Ile	Pro	Asp	
				305					310					315	
Tyr	Met	Pro	Leu	Trp	Val	Tyr	Trp	Phe	Ile	Thr	Gly	Ile	Ser	Ile	
				320					325					330	
Leu	Leu	Val	Gly	Ser	Val	Ile	Leu	Leu	Ile	Val	Cys	Met	Thr	Trp	
				335					340					345	
Arg	Leu	Ala	Gly	Pro	Gly	Ser	Glu	Lys	Tyr	Ser	Asp	Asp	Thr	Lys	
				350					355					360	

Tyr Thr Asp Gly	Leu	Pro	Ala	Ala	Asp	Leu	Ile	Pro	Pro	Pro	Leu	
	365					370					375	
Lys Pro Arg Lys	Val	Trp	Ile	Ile	Tyr	Ser	Ala	Asp	His	Pro	Leu	
	380					385					390	
Tyr Val Asp Val	Val	Leu	Lys	Phe	Ala	Gln	Phe	Leu	Leu	Thr	Ala	
	395					400					405	
Cys Gly Thr Glu	Val	Ala	Leu	Asp	Leu	Leu	Glu	Glu	Gln	Ala	Ile	
	410					415					420	
Ser Glu Ala Gly	Val	Met	Thr	Trp	Val	Gly	Arg	Gln	Lys	Gln	Glu	
	425					430					435	
Met Val Glu Ser	Asn	Ser	Lys	Ile	Ile	Val	Leu	Cys	Ser	Arg	Gly	
	440					445					450	
Thr Arg Ala Lys	Trp	Gln	Ala	Leu	Leu	Gly	Arg	Gly	Ala	Pro	Val	
	455					460					465	
Arg Leu Arg Cys	Asp	His	Gly	Lys	Pro	Val	Gly	Asp	Leu	Phe	Thr	
	470					475					480	
Ala Ala Met Asn	Met	Ile	Leu	Pro	Asp	Phe	Lys	Arg	Pro	Ala	Cys	
	485					490					495	
Phe Gly Thr Tyr	Val	Val	Cys	Tyr	Phe	Ser	Glu	Val	Ser	Cys	Asp	
	500					505					510	
Gly Asp Val Pro	Asp	Leu	Phe	Gly	Ala	Ala	Pro	Arg	Tyr	Pro	Leu	
	515					520					525	
Met Asp Arg Phe	Glu	Glu	Val	Tyr	Phe	Arg	Ile	Gln	Asp	Leu	Glu	
	530					535					540	
Met Phe Gln Pro	Gly	Arg	Met	His	Arg	Val	Gly	Glu	Leu	Ser	Gly	
	545					550					555	
Asp Asn Tyr Leu	Arg	Ser	Pro	Gly	Gly	Arg	Gln	Leu	Arg	Ala	Ala	
	560					565					570	
Leu Asp Arg Phe	Arg	Asp	Trp	Gln	Val	Arg	Cys	Pro	Asp	Trp	Phe	
	575					580					585	
Glu Cys Glu Asn	Leu	Tyr	Ser	Ala	Asp	Asp	Gln	Asp	Ala	Pro	Ser	
	590					595					600	
Leu Asp Glu Glu	Val	Phe	Glu	Glu	Pro	Leu	Leu	Pro	Pro	Gly	Thr	
	605					610					615	
Gly Ile Val Lys	Arg	Ala	Pro	Leu	Val	Arg	Glu	Pro	Gly	Ser	Gln	
	620					625					630	
Ala Cys Leu Ala	Ile	Asp	Pro	Leu	Val	Gly	Glu	Glu	Gly	Gly	Ala	
	635					640					645	
Ala Val Ala Lys	Leu	Glu	Pro	His	Leu	Gln	Pro	Arg	Gly	Gln	Pro	
	650					655					660	
Ala Pro Gln Pro	Leu	His	Thr	Leu	Val	Leu	Ala	Ala	Glu	Glu	Gly	
	665					670					675	
Ala Leu Val Ala	Ala	Val	Glu	Pro	Gly	Pro	Leu	Ala	Asp	Gly	Ala	
	680					685					690	
Ala Val Arg Leu	Ala	Leu	Ala	Gly	Glu	Gly	Glu	Ala	Cys	Pro	Leu	
	695					700					705	
Leu Gly Ser Pro	Gly	Ala	Gly	Arg	Asn	Ser	Val	Leu	Phe	Leu	Pro	
	710					715					720	
Val Asp Pro Glu	Asp	Ser	Pro	Leu	Gly	Ser	Ser	Thr	Pro	Met	Ala	
	725					730					735	
Ser Pro Asp Leu	Leu	Pro	Glu	Asp	Val	Arg	Glu	His	Leu	Glu	Gly	
	740					745					750	

Leu	Met	Leu	Ser	Leu	Phe	Glu	Gln	Ser	Leu	Ser	Cys	Gln	Ala	Gln
				755					760					765
Gly	Gly	Cys	Ser	Arg	Pro	Ala	Met	Val	Leu	Thr	Asp	Pro	His	Thr
				770					775					780
Pro	Tyr	Glu	Glu	Glu	Gln	Arg	Gln	Ser	Val	Gln	Ser	Asp	Gln	Gly
				785					790					795
Tyr	Ile	Ser	Arg	Ser	Ser	Pro	Gln	Pro	Pro	Glu	Gly	Leu	Thr	Glu
				800					805					810
Met	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Gln	Asp	Pro	Gly	Lys	Pro	Ala
				815					820					825
Leu	Pro	Leu	Ser	Pro	Glu	Asp	Leu	Glu	Ser	Leu	Pro	Ser	Leu	Gln
				830					835					840
Arg	Gln	Leu	Leu	Phe	Arg	Gln	Leu	Gln	Lys	Asn	Ser	Gly	Trp	Asp
				845					850					855
Thr	Met	Gly	Ser	Glu	Ser	Glu	Gly	Pro	Ser	Ala				
				860					865					

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